

Supplemental Information

Figure legends

Figure 1

(A) Cell growth activity measured by MTT assay showed similar results to the results obtained by luciferase assay. After the 3-days incubation of CM derived from PNT-2 cells or PC-3M-luc cells with PC-3M cells, inhibitory effect of PC-3M-luc cellular proliferation was assessed by MTT assay and luciferase assay. Each bar is presented as mean S.E. (n=3). (**, $p<0.005$; Student's *t* test) (B) Purified exosomes secreted from PNT-2 cells are enriched in CD63 protein. The exosome fractions collected from PNT-2 derived CM incubated for 0 hr or 48 hr and whole cell lysate of PNT-2 cells were analyzed by immunoblotting with an anti-CD63 antibody or anti- β -actin antibody. Equal amounts of proteins from exosome and whole cells were analyzed. Twenty microlitter of exosome samples derived from 0 h incubation and 48 h were also analyzed. CD63 is heavily glycosylated displaying a range of bands between 50 and 75 kDa. (C) After the treatment with the 10 μ M of GW4869 for 48 hr, the total amounts of proteins in the exosomal pellet purified from large scale cultures of PNT-2 cells were quantified by a BCA assay and are presented as the values per 10 million secreting cells. Each bar is presented as mean S.E. (n=3). (**, $p<0.005$; Student's *t* test) (D) Fluorescence photos of PC-3M-luc cells incorporating PKH67-loaded exosomes isolated from PNT-2 cells. PC-3M-luc cells were incubated for 0 hr and 16 hr with PKH67-loaded exosomes. Emission at 514 nm was detected with Eclipse TE 2000 Inverted Research Microscope and images were produced by NIS-Elements BR software. The size bar indicates 50 μ m.

Figure 2

(A) GW4869 treatment of PNT-2 did not affect the expression of miR-143 in the cells. PNT-2 cells were treated with 10 μ M of GW4869 in a 6-well plate. One day later, medium were changed to non-serum medium. The following day, cell extract were applied to QRT-PCR for miR-143. The values on the *y* axis are depicted relative to the amount of miRNAs of control, which is arbitrarily defined as 1. (B) siRNA-mediated knockdown of nSMase2 in PNT-2 cells. PNT-2 cells were transfected with either negative control or nSMase 2 siRNA in a 24-well plate. One day later, cell extract were applied to QRT-PCR for nSMase 2. The values on the *y* axis are depicted relative to the amount of nSMase 2 of control, which is arbitrarily defined as 1. (**, $p<0.005$; Student's *t* test) (C) and (D) siRNA-mediated knockdown of nSMase 2. PNT-2 cells

were transfected with either negative control or nSMase2 siRNA in a 24-well plate. The following day, conditioned medium (C) and cell extract (D) were applied to QRT-PCR for miR-143. The values on the *y* axis are depicted relative to the amount of miRNAs of control, which is arbitrarily defined as 1. (**, $p < 0.005$; Student's *t* test)

Figure 3

(A) miR-143 were secreted into culture medium. Conditioned medium from miR-143 stably expressing HEK293 cells or control non-transfected HEK293 cells were harvested and performed QRT-PCR as described under Materials and methods. The values on Y-axis are depicted relative to the amount of miR-143 in conditioned medium from non-transfected HEK293 cells, which is arbitrarily defined as 1. (B) PC-3M-luc cells were incubated with CM derived from HEK293 cells or over-producing miR-143 HEK293 cells for 24 hr at 37°C and the transfer of miR-143 was evaluated by qRT-PCR. The values on the *y* axis are depicted relative to the normalized miR-143 expression of original HEK293 CM-treated cells, which is defined as 1. Each *bar* is presented as the mean S.E. ($n=3$). (*, $p < 0.05$; Student's *t* test)

Fig. 1

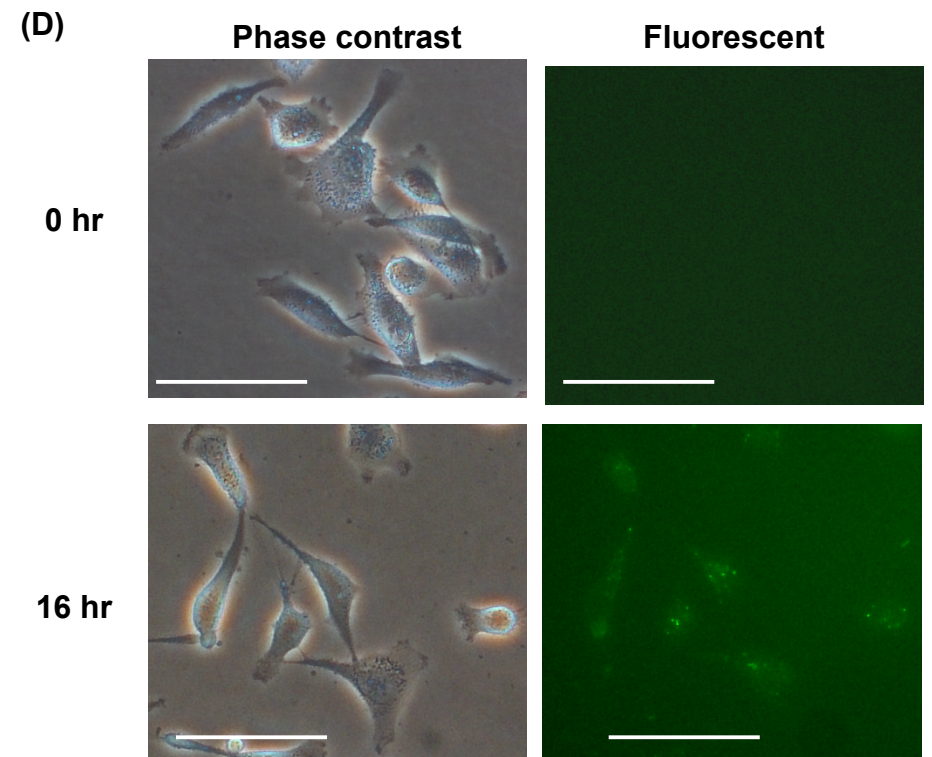
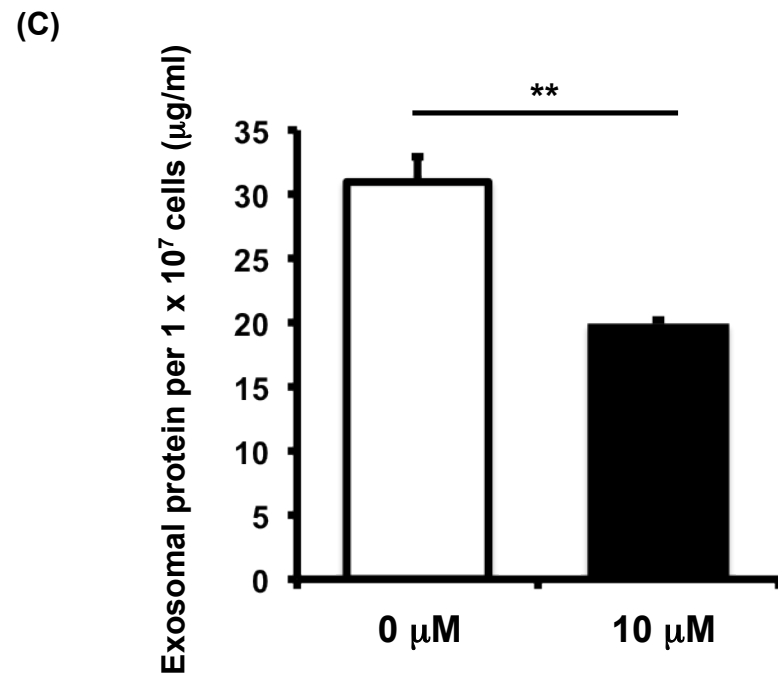
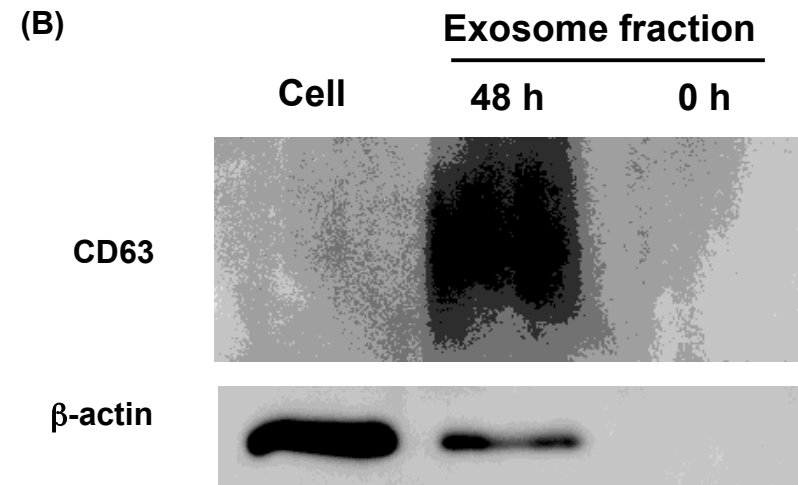
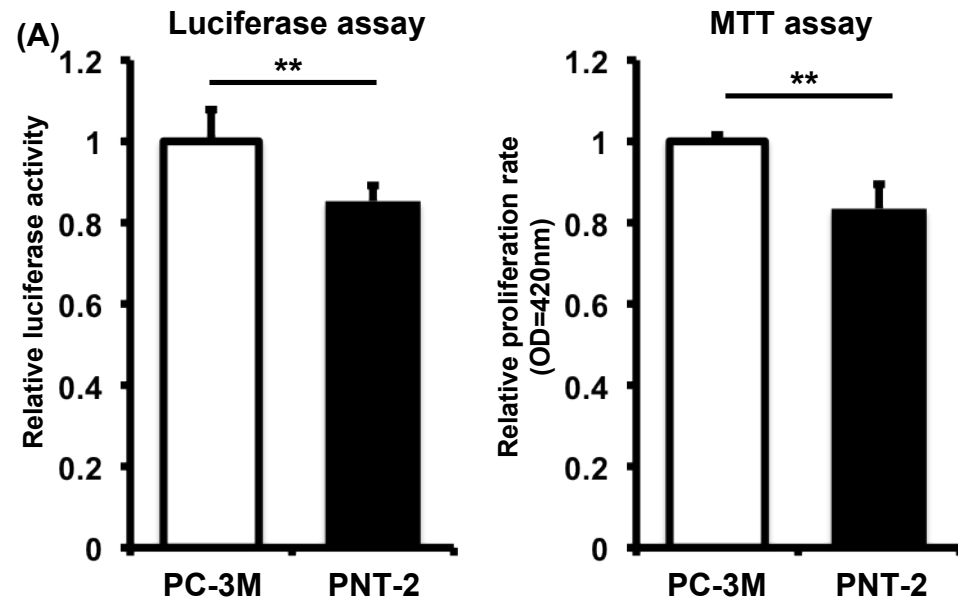
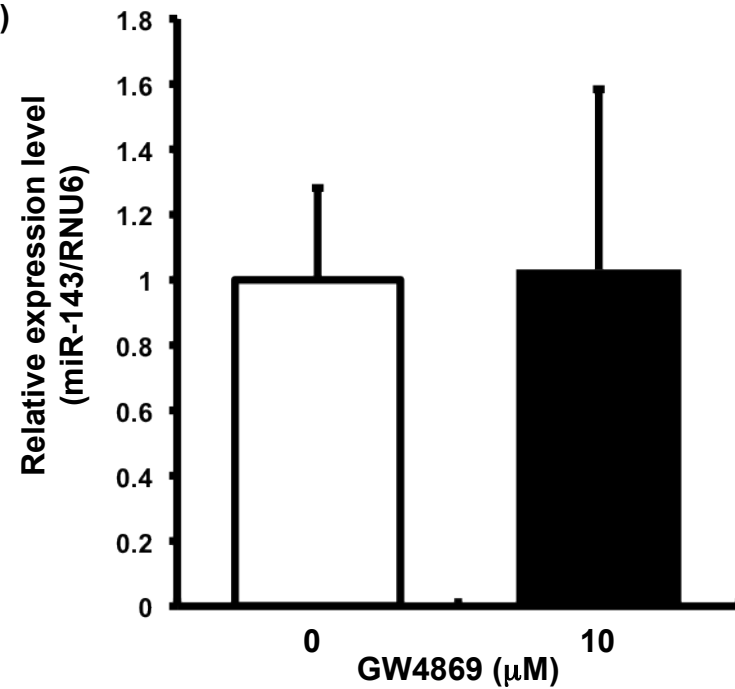
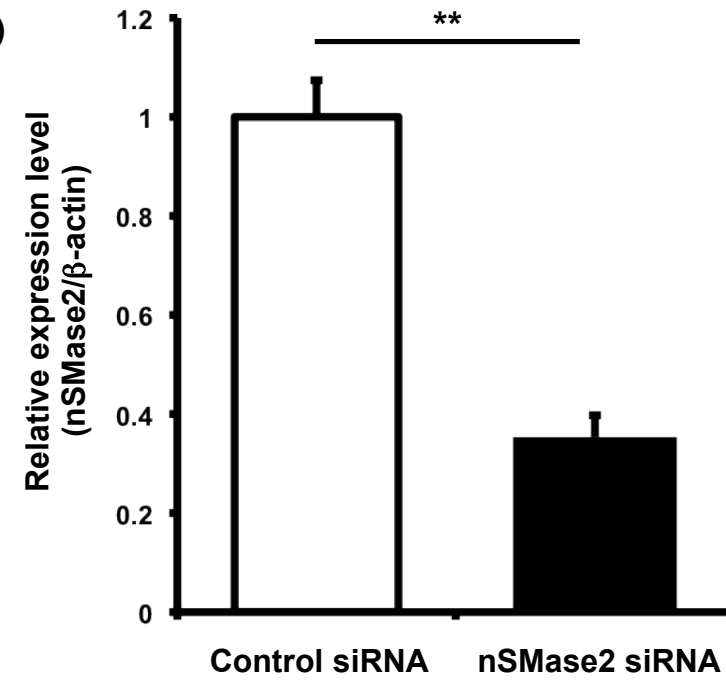


Fig. 2

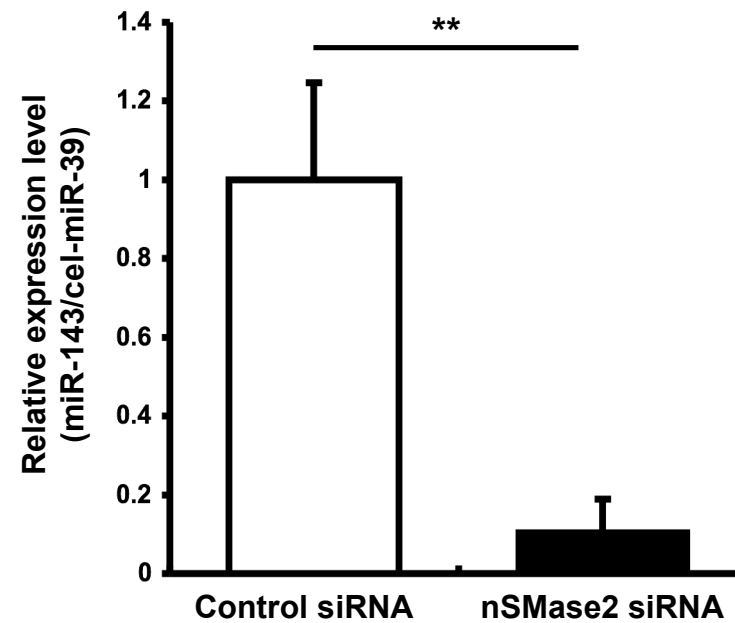
(A)



(B)



(C)



(D)

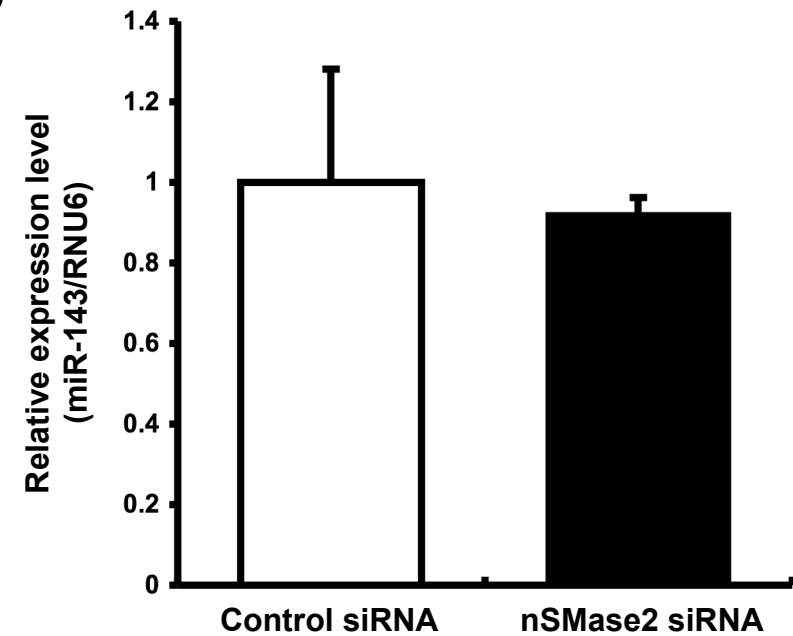


Fig. 3

